

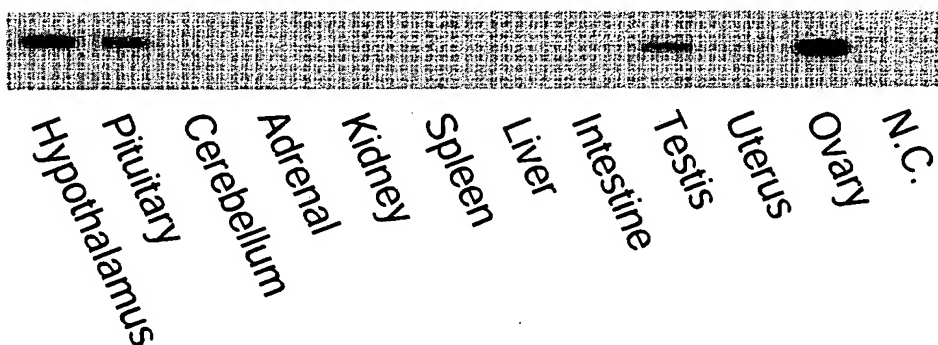
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Fig. 1

1 MSDGNPELLSTSQTYN SQGESNEDYEIPPITPPNLPEPSLLHLGDHEAGY 50  
51 HSLCHGLAPNGLLPAYSYQAMDLP AIMVSNMLAQDGHLLSGQLPTIQEMV 100  
101 HSEVAAYDSGRPGPLLGRPAMLASHMSALSQSQLISQMGLRSGIAHSSPS 150  
151 PPGSKSATPSPSSSTQEEESDAHFKISGEKRPSTDPGKKAKNPKKKKKKD 200  
201 PNEPQKPVSA YALFFRDTQAAIKGQNPSATFGDVSKI VASMWDSLGE EQK 250  
251 QAYKRKTEAAKKEYLKALAA YRASLVSKSPPDQGEAKNAQANPPAKMLPP 300  
301 KQPMYAMPGLASFLTPSDLQAFRSAASPASLARTLGSKALLPGLSTSPPP 350  
351 PSFPLSPSLHQQLPLPPHAQG TLLSPPLSMSPAPQPPVLPAPMALQVQLA 400  
401 MSPSPPGPQDFPHISDFPSGSGSRSPGPSNPSSSGDWDG SYPSGERGLGT 450  
451 CRLCRSSPPPTTSPKNLQEPSAR 473

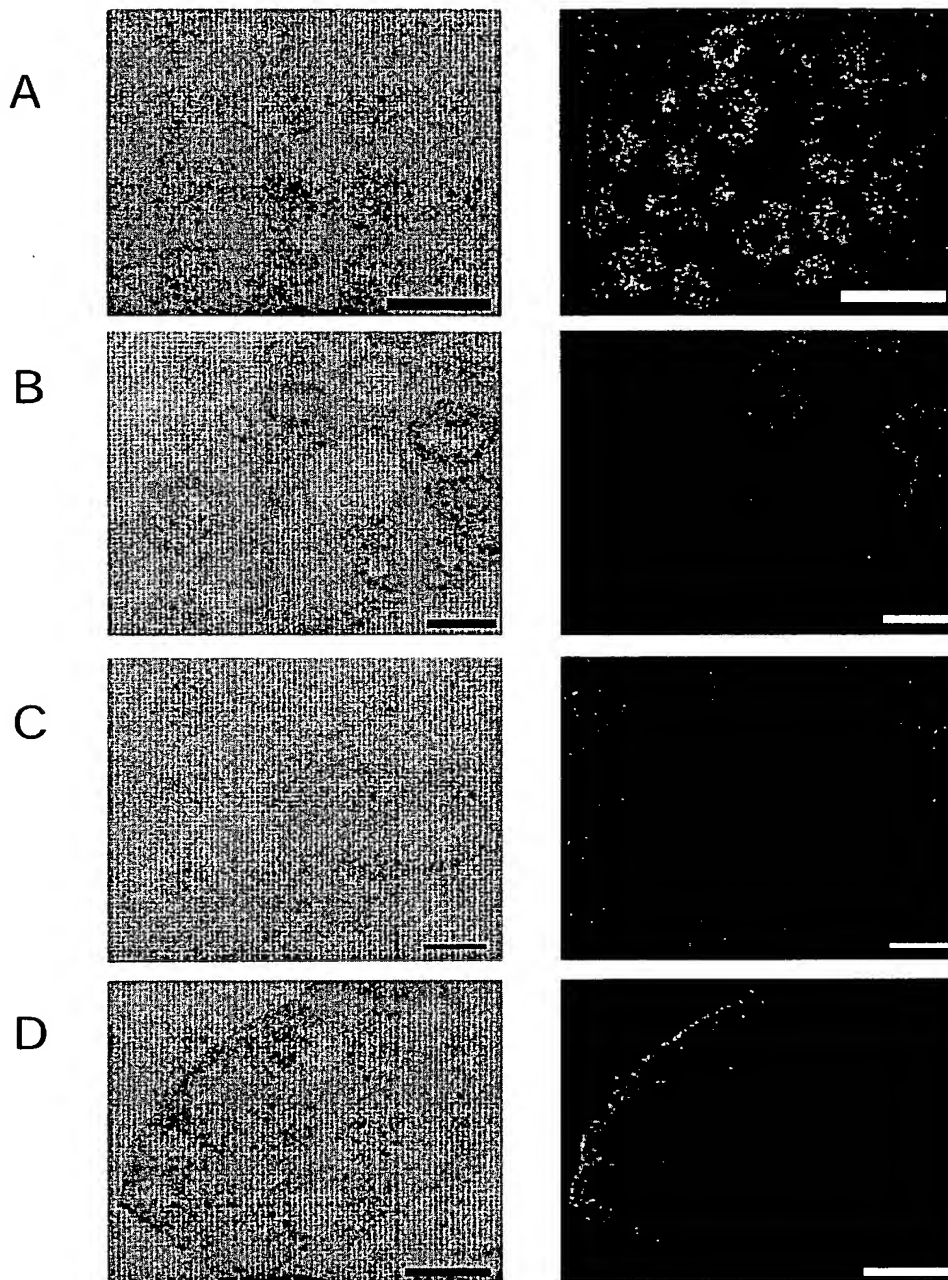
Fig. 2

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Fig. 3



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Fig. 4

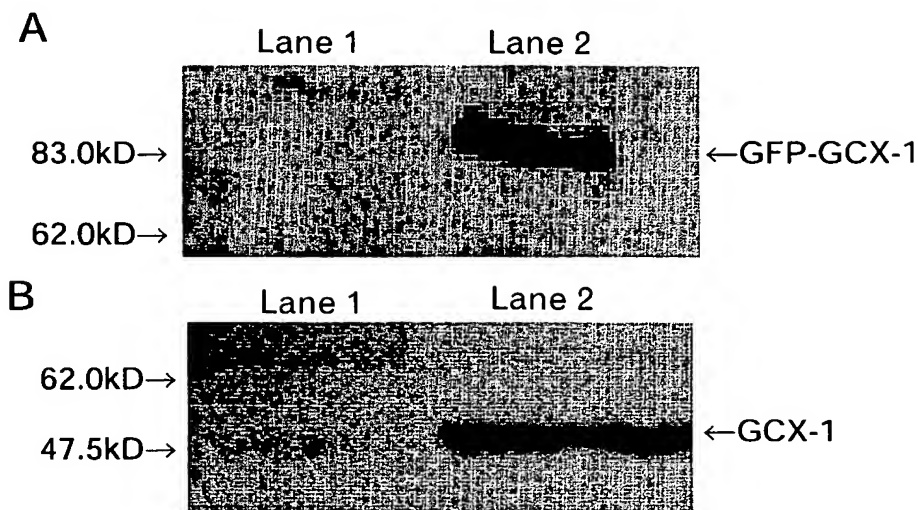
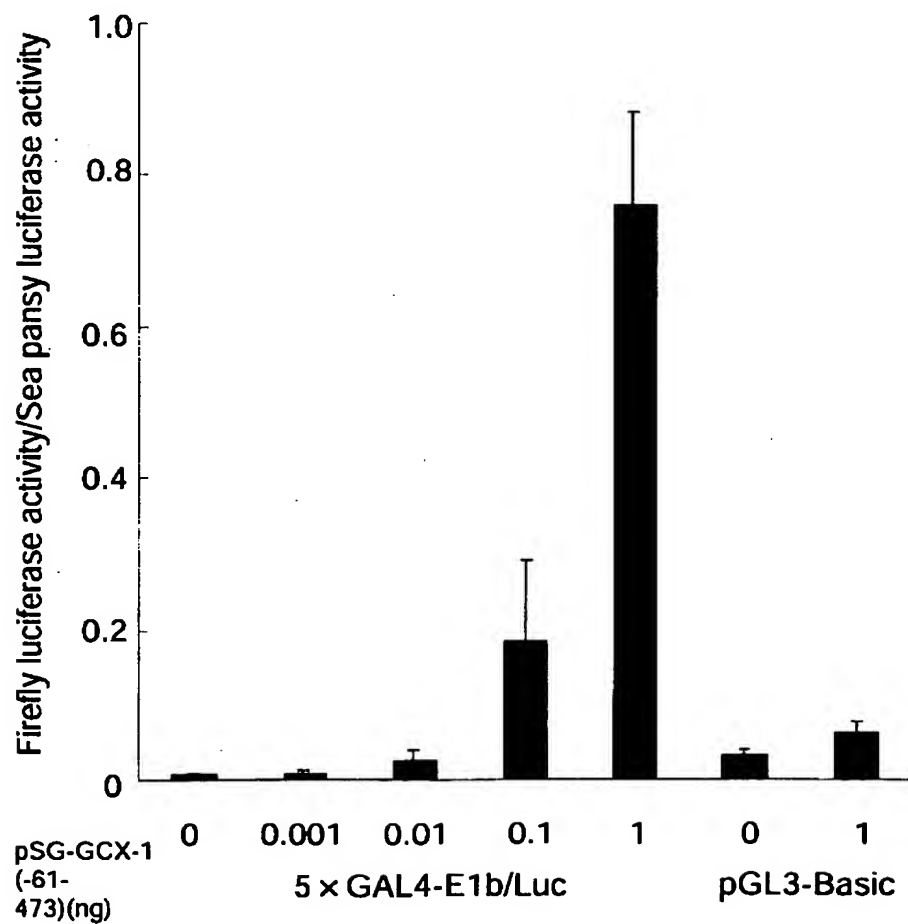


Fig. 5



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Fig. 6

